	Changed a file from non-ASCII to ASCII
	Changed the margins in cases where the sequence text was "wrapped," down to the next line.
	Edited a format error in the Current Application Data section, specifically:
•	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII *garbage* at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
-	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted ending stop codon in amino acid sequences and adjusted the *(A)Length: field accordingly (error life to a Patentin bug). Sequences corrected:
	Other: Segr 2, 7, 13-20 - isserted fard returns

#Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

1646

Does Not Comply RAW SEQUENCE LISTING DATE: 12/26/2000 Corrected Diskette Needed PATENT APPLICATION: US/09/589,777A TIME: 12:57:35 Input Set : A: \14401023011, txt Output Set: N:\CRF3\12262000\1589777A.raw 4 <110> APPLICANT: Sukhatme, Vikas P. b <120> TITLE OF INVENTION: Anti-Angiogenic Peptides and Methods of Use Thereof 9 <130> FILE REFERENCE: 1440.1023-011 11 <140> CURRENT APPLICATION NUMBER: US 09/589,777A. 12 <141> CURRENT FILING DATE: 2000-06-08 14 <150> PRIOR APPLICATION NUMBER: PCT/US98/26057 15 <151> PRIOR FILING DATE: 1998-11-16 17 <150> PRIOR APPLICATION NUMBER: US 60/108,536 BEST AVAILABLE COPY 18 <151> PRTOR FILING DATE: 1998-04-22 20 <150> PRIOR APPLICATION NUMBER: US 60/082,663 21 <151> PRIOR FILING DATE: 1998-04-22 23 <150> PRIOR APPLICATION NUMBER: US 60/067,888 24 <151> PRIOR FILING DATE: 1997-12-07 26 <160> NUMBER OF SEQ ID NOS: 23 28 <170> SOFTWARE: FastSEQ for Windows Version 4.0 ERRORED SEQUENCES 97 <210> SEQ ID NO: 2 98 <211> LENGTH: 184 99 <212> TYPE: PRT 100 <213> ORGANISM: Mus musculus 102 <400> SEQUENCE: 2 103 His Thr His Gln Asp Phe Gln Pro Val Leu His Leu Val Ala Leu Asn 1.0 105 Thr Pro Leu Ser GI; Gly Met Arg Gly IIe Arg Gly Ala Asp Phe GIn 106 $^{\circ}$. 20 $^{\circ}$ 25 $^{\circ}$ 30 107 Cys Phe Gln Gln Ala Arg Ala Val Gly Len Ser Gly Thr Phe Arg Ala 35 40 109 Phe Leu Ser Ser Arg Leu Gln Asp Leu Tyr Ser Ile Val Arg Arg Ala 110 50 55 60

1.11 Asp Arg Gly Ser Val Pro 11e Val Asn Leu Lys Asp Glu Val Leu Ser 112 65 70 75 80

113 Pro Ser Trp Asp Ser Leu Phe Ser Gly Ser Gin Gly Gln Leu Gln Pro

115 Gly Ala Arg Ile Phe Ser Phe Asp Gly Arg Asp Val Leu Arg His Pro 116 \$100\$117 Ata Trp Pro Gli Lys Ser Val Trp His Gly Ser Asp Pro Ser Gly Arg

.120 119 Arg Leu Met Glu Ser Tyr Cys Glu Thr Trp Arg Thr Glu Thr Thr Gly 120 $^{\circ}$ 130 $^{\circ}$ 135 $^{\circ}$ 140

121 Ala Thr Gly Gln Ala Ser Ser Leu Leu Ser Gly Arg Leu Leu Glu Gln

70

150

Lys Ala Ala Ser Cys His Asn Ser Tyr Ile Val Leu Cys Ile Glu Asn

8.5

118 115

146 <210> SEQ ID NO: 5

E--> 123

, usert hard return

Input Set : A:\14401023011.txt Output Set: N:\CRF3\12262000\1589777A.raw 147 <211> LENGTH: 24 148 <212> PYP': PRT 149 <213> OPGANTSH: Artificial Sequence 151 <220> FERTURES 152 <223> OTHER INFORMATION: Leader peptide on protein produced by prokaryotic 153 expression system pET17b, mouse endostatin begins immediately after. . 154 156 <400> SEQUENCE: 5 E--> 1.57 Met Gly His His His His His His His His His Ser Ser Gly His 1 10 170 <210> SEQ ID NC: 7 171 <211> LENGTH: 21 172 <212> TYPE: PRT 173 <213> ORGANISM: Artificial Sequence 175 <220> PEATURE: 176 <223> OTHER INFORMATION: Leader peptide on protein produced by prokaryotic 177 $\exp \operatorname{ression}$ system pET28a, mouse endostatin begins immediately after. 178 180 <400> SEQUENCE: 7 E--> 181 Met Gly Ser Ser His His His His His Ser Ser Gly Leu Val Pro 1 10 237 <210> SEQ ID NO: 13 238 <211> LENGTH: 26 239 <212> TYPE: PRT 240 <213> ORGANTSM: Artificial Sequence 242 <220> FEATURE: 243 <223> OTHER INFORMATION: Leader peptide on protein produced by eukaryotic

DATE: 12/26/2000

TIME: 12:57:35

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/589,777A

yeast expression system pPTCZaA, mouse endostatio

yeast expression system pPICZaA, mouse endostatin

321 <223> OTHER INFORMATION: Leader peptide on protein produced by eukaryotic

5

protein begins immediately after.

Glu Phe Met Gly His His His His His His His His His Ser Ser l

protein begins immediately after.

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244

245

322

323

E--> 248

247 <400> SEQUENCE: 13

315 <210> SEQ ID NO: 20 316 <211> LENGTH: 8 317 <21.2> TYPE: PRT

325 <400> SEQUENCE: 20

E--> 326 Glu Phe His His His His His Ti

320 <220> FEATURE:

318 <213> ORGANISM: Artificial Sequence

10

VERIFICATION SUMMARY PATENT APPLICATION: US/09/589,777A

DATE: 12/26/2000 TIME: 12:57:37

Input Set : A:\14401023011.txt
Output Set: N:\CRF3\12262000\1589777A.raw

L:123 M:252 E: No. of Seq. differs, <211>LENGTH:Input:184 Found:160 SEQ:2 L:157 M:252 E: No. of Seq. differs, <211>LENGTH:Input:24 Found:0 SEQ:5 L:181 M:252 E: No. of Seq. differs, <211>LENGTH:Input:21 Found:0 SEQ:7 L:248 M:252 E: No. of Seq. differs, <211>LENGTH:Input:26 Found:0 SEQ:13 L:326 M:252 E: No. of Seq. differs, <211>LENGTH:Input:8 Found:0 SEQ:20

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